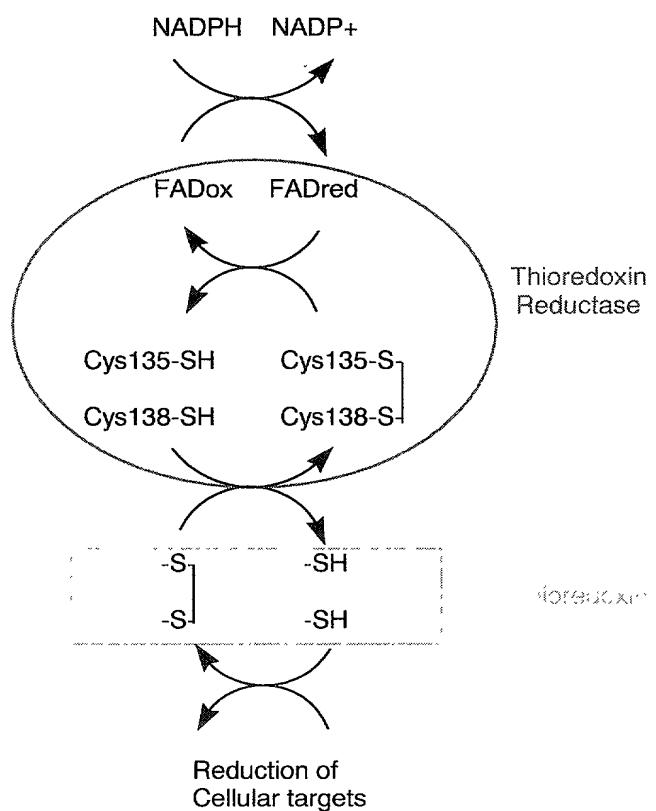


Figure 1



Title: CRYSTALLIZATION AND STRUCTURE DETERMINATION OF

STAPHYLOCOCCUS AUREUS THIOREDOXIN REDUCTASE

Applicant(s): Timothy E. BENSON **Serial No.:** Unassigned
P.O. Box 88822 U.S.C. F. M. N.Y. F.I. 77-1888822 U.S.

Docket: 00032.US1 Express Mail No.: EL 776900927 US

Filed: Herewith

Sheet 2 of 13

Figure 2

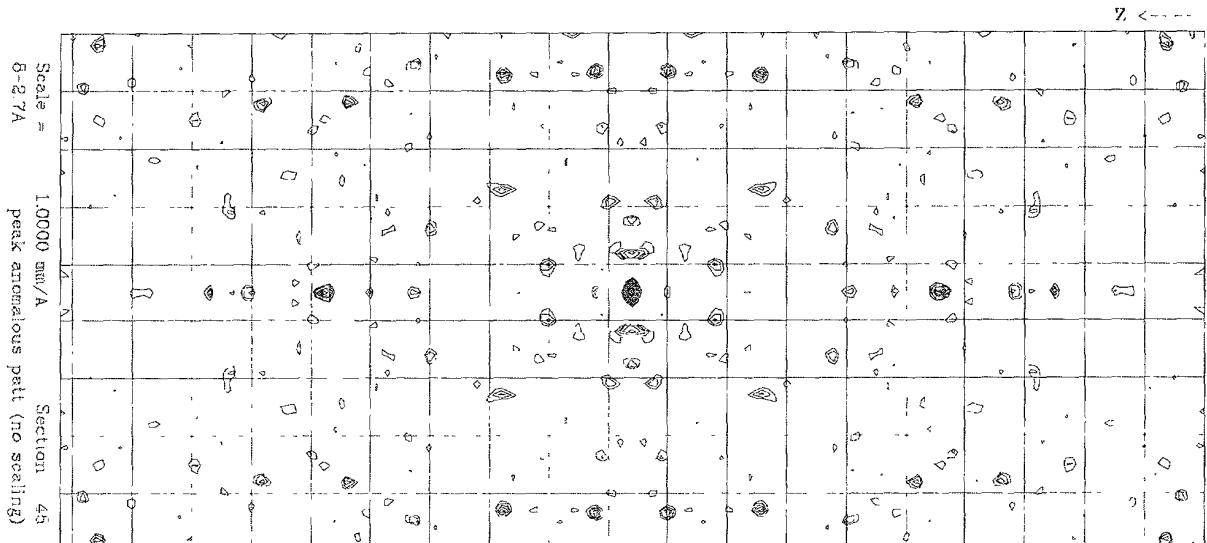
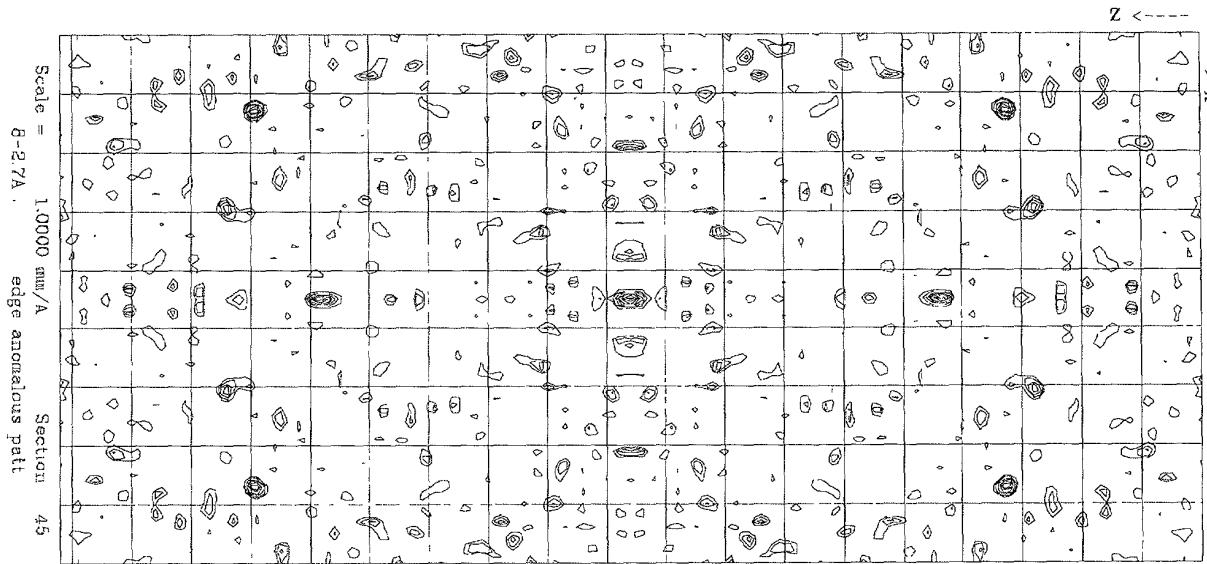
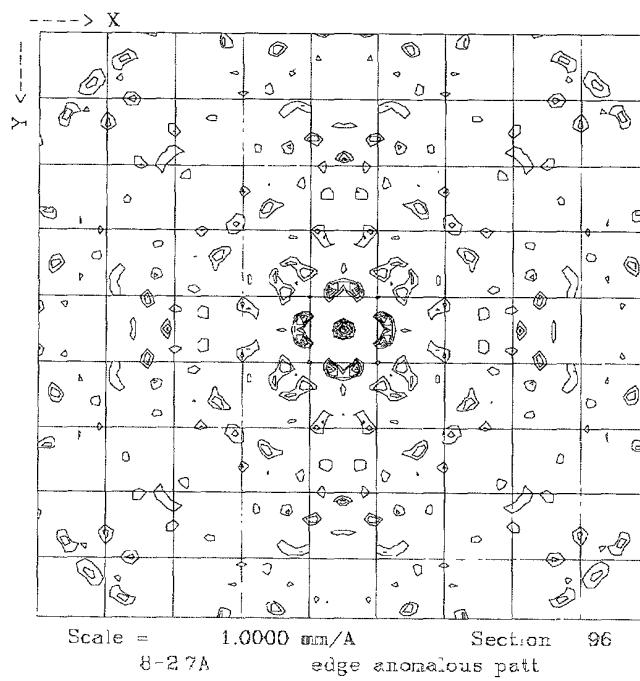
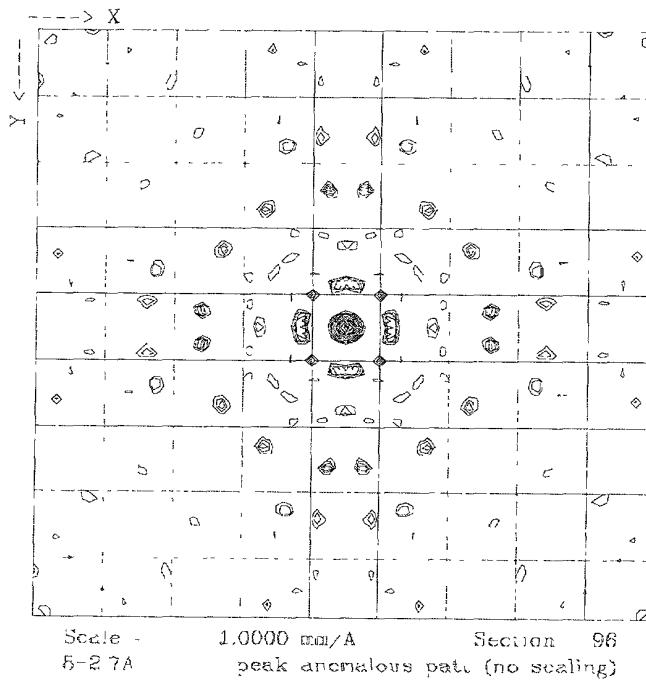


Figure 3

a.



b.



Title: CRYSTALLIZATION AND STRUCTURE DETERMINATION OF

STAPHYLOCOCCUS AUREUS THIOREDOXIN REDUCTASE

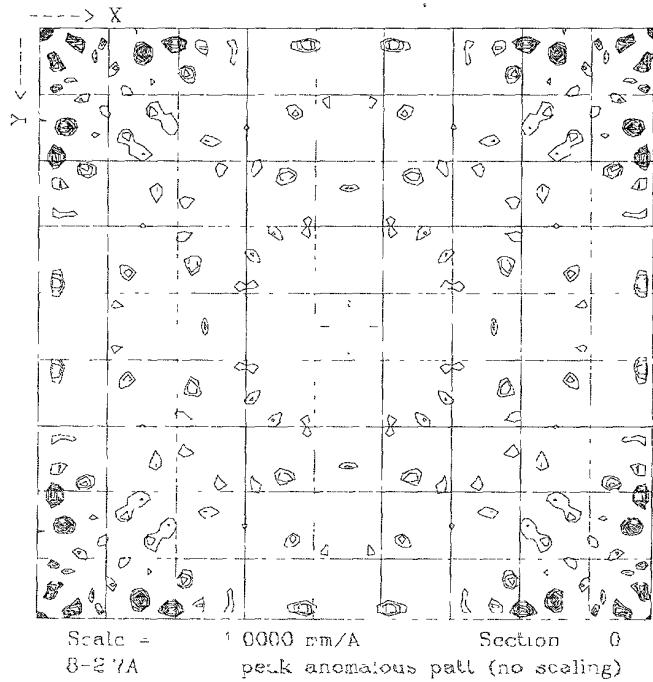
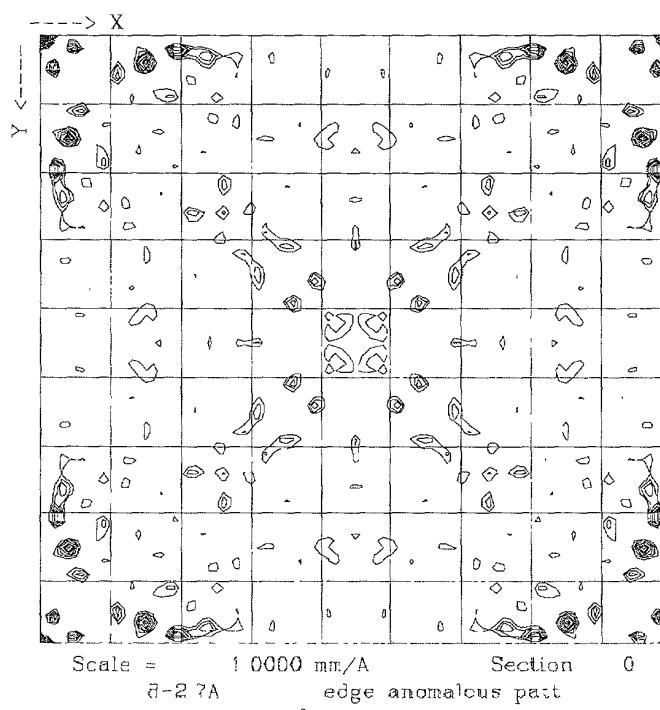
Applicant(s): Timothy E. BENSON Serial No.: Unassigned

Docket: 00032.US1 Express Mail No.: EL 776900927 US

Filed: Herewith

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Figure 4



Title: CRYSTALLIZATION AND STRUCTURE DETERMINATION OF

STAPHYLOCOCCUS AUREUS THIOREDOXIN REDUCTASE

Applicant(s): Timothy E. BENSON **Serial No.:** Unassigned
U.S. PATENT AND TRADEMARK OFFICE

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Figure 5

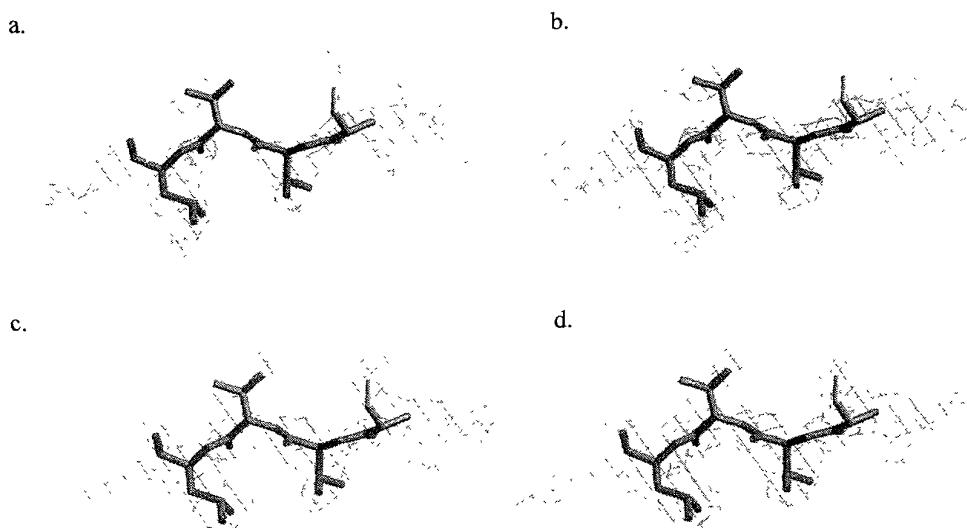


Figure 6

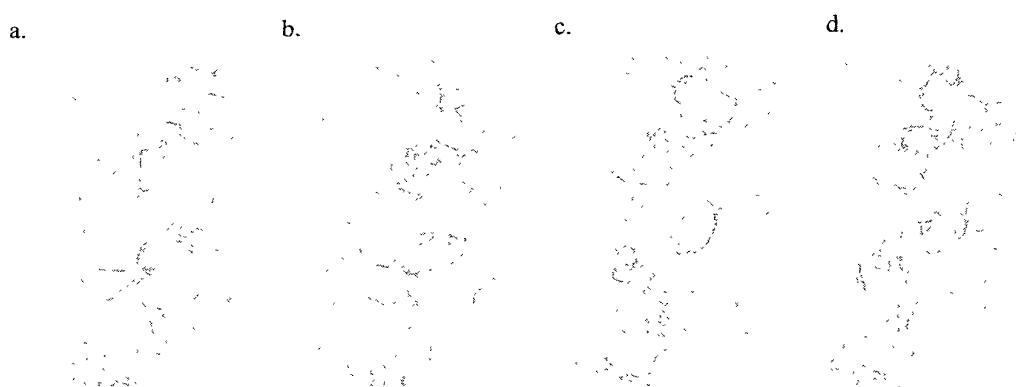


Figure 7

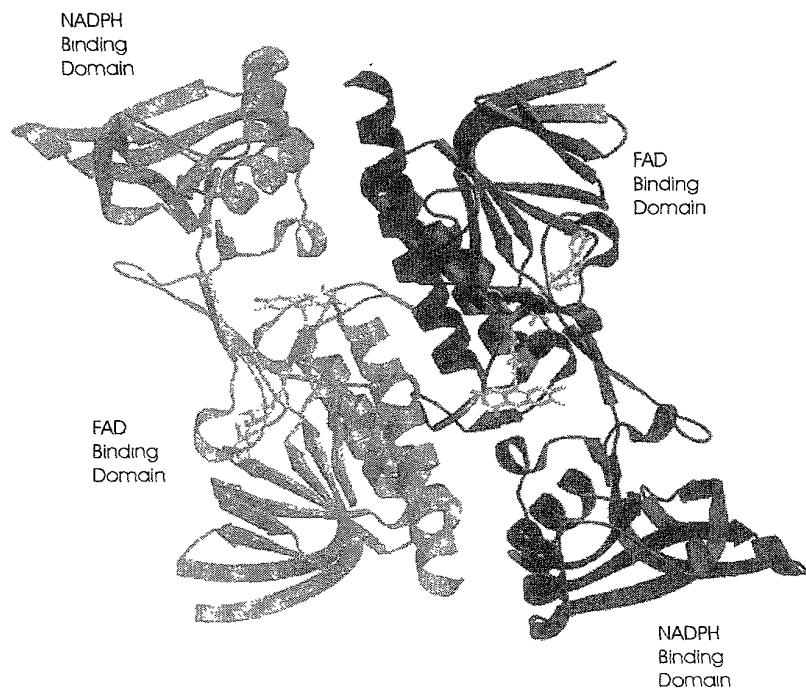
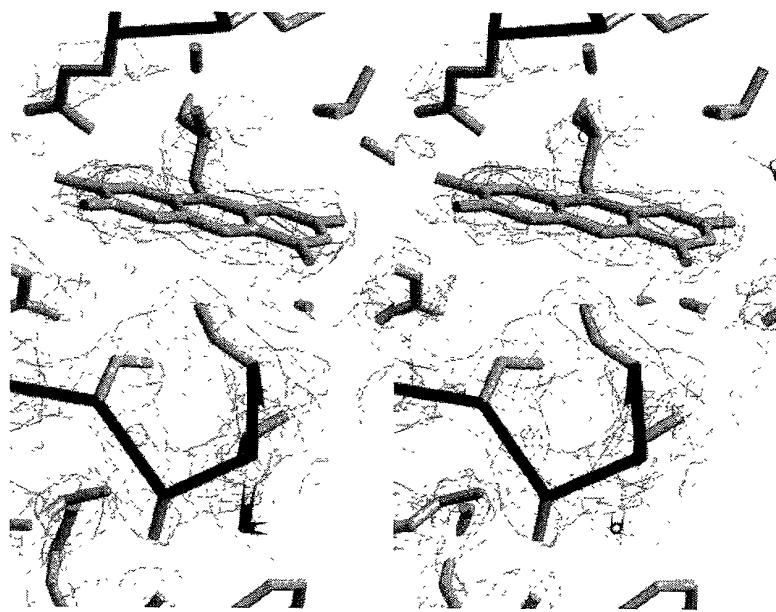


Figure 8

a.



b.

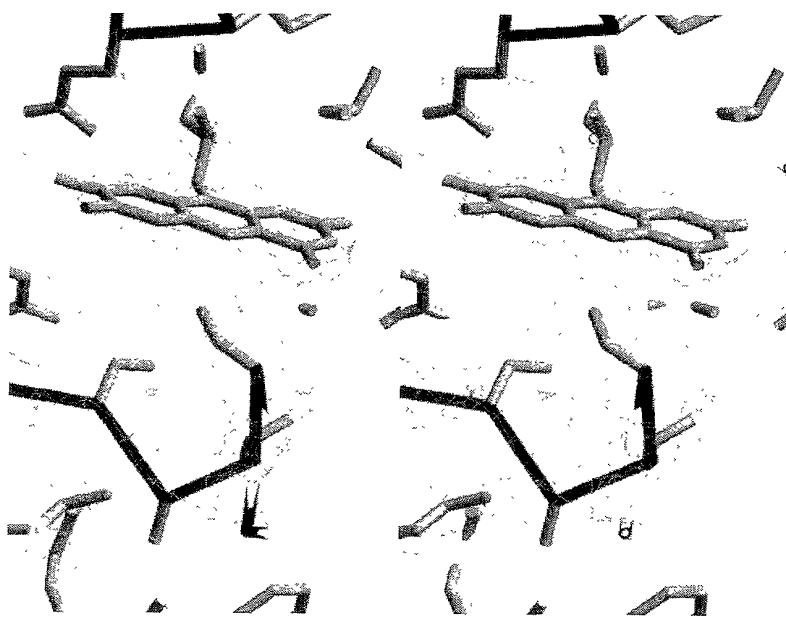
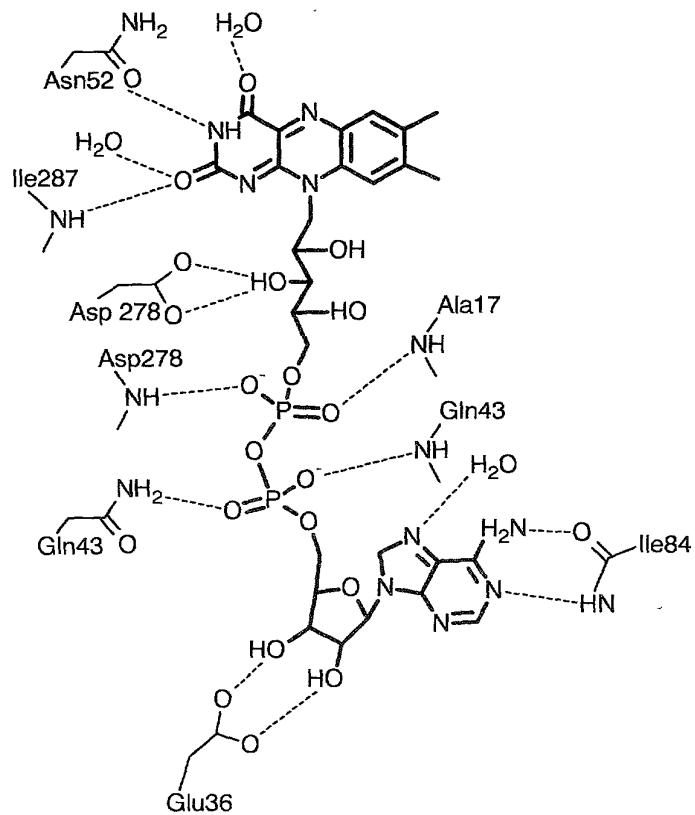


Figure 9



Filed: Herewith
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Figure 10

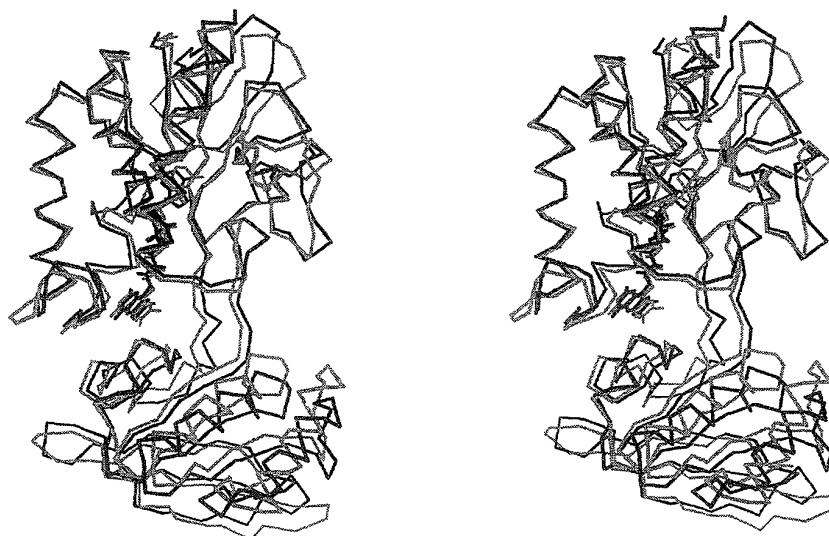


Figure 11

(SEQ ID NO: 2)	TrxB_ecoli	1	500
(SEQ ID NO: 1)	TrxB_sa	(1) -GTTKHSKILLIGSGPAGYAAVYAAARANLQPVEITGMEEKGGQIHTTTEV	
		(1) MGTEIDFDIAITGAGPAGMTAAVYASRANLKTVMIERGIPGGQMANTEEV	
		51	100
TrxB_ecoli	(50)	ENWP GDPN DLTGP PLLMER MHEHATKFETE IIFDHINKV DLQNRPFRLNGD	
TrxB_sa	(51)	ENFP G-FEM ITGP DLST KMFE HAKK FGAV YQY CDIK SVED KGEY KV INFG	
		101	150
TrxB_ecoli	(100)	NGE YTCD AHI IATG ASARY LGL PSEE AFK GRGV SACAT CDG FFYRNOKVA	
TrxB_sa	(100)	NKEL TAKA VII IATGA EYKK TGV PG EQL GG RGV SYCAV CDGA FFKN KRLF	
		151	200
TrxB_ecoli	(150)	VIGGG NTAVEE ALV LSN IA SEV HLTH RR DG FRAE KIL IKR LM DKV ENGN I	
TrxB_sa	(150)	VIGGG DSA VE EGTF TTKF ADKV TIV HRR DEL RA QRI LQDRA FK N--DKI	
		201	250
TrxB_ecoli	(200)	ILHT NR TTEE VTGD QM GV TL RDT QNS DNIE SLDV AGLF VAIGH SPNT	
TrxB_sa	(197)	DFI WS HTT KSIN EKD GKV GS VTL ST-KDG SEET HADGV FTY IGM KPLT	
		251	300
TrxB_ecoli	(250)	AIFEGQ -LELE NGYIKV QOSGI HG NAT QT S I PGV FAAG DVMD HIY RQ AITS A	
TrxB_sa	(246)	APFK DL GIT NDVG YIV TK--DD---MTT SVPG IFAAG DV RD KGL RQIV TAT	
		301	329
TrxB_ecoli	(300)	GTGC MAAL DAERYL DGL ADAK-----	
TrxB_sa	(292)	GDGS TAAQS AAEY IEHL NDO ARSHHHHHH	

Figure 12

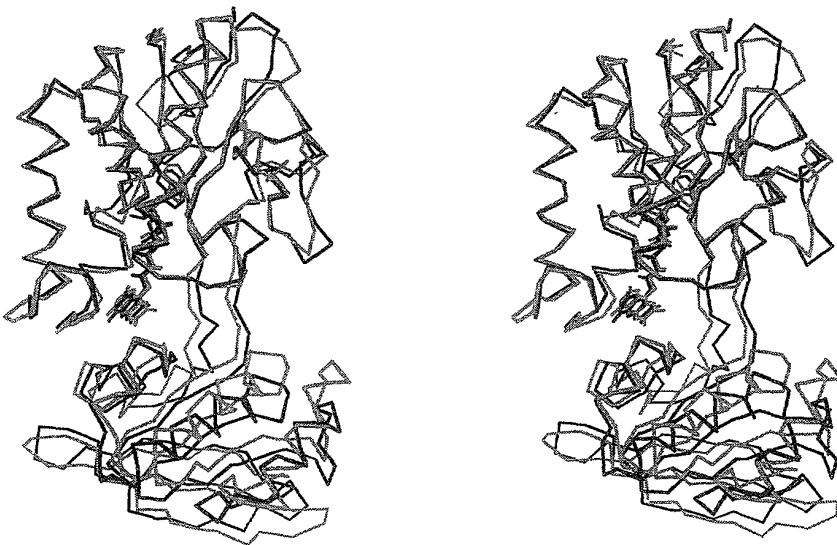


Figure 13

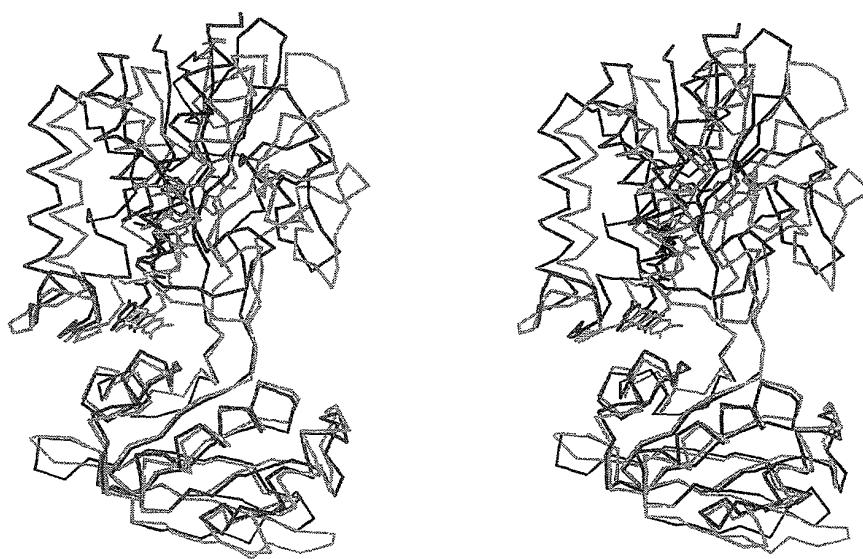


Figure 14

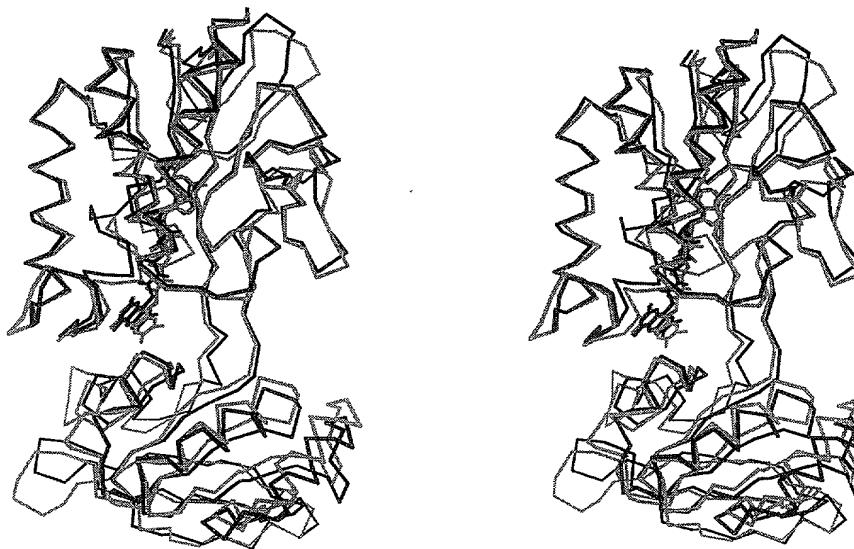


Figure 15

(SEQ ID NO: 3)	Tr_athaliana	1	50
(SEQ ID NO: 1)	TrxB_sa	(1) MNGLETHNTRLCIVGSGPAAHAAIYARAEELKPLIFEGWMANDIAPGGQ (1) -MGTEID-FDTAIIAGPAGMTAAVYASRANLKTVMIERG----IPGGQ	
		51	100
Tr_athaliana	(51) LTTTDPVENFPGFPEGILGVELTDKFRKQSERFGTTIFTETVTKVDFSSK		
TrxB_sa	(44) MANTEEVENFPGF-EMITGPDLSTKMFEHAKKFGAVYQYGDIKSVEDKGK		
		101	150
Tr_athaliana	(101) PFKLFTDSKAILADAVILAIGAVAKRISFVGSGEVLGGFWNRGTSACAVC		
TrxB_sa	(93) YKVINFGNKEELTAKAVIILATGAEYKKIGVPGEQEYGG---RGYSYCAVC		
		151	200
Tr_athaliana	(151) DGAAPIFRNKPLAVIGGGDSAMEEANFLTKYGSKVYIITHRRDAFRASKIM		
TrxB_sa	(139) DGA--FFKNKRLFVIGGGDSAVEEGTFLTKFADKVTVHRRDELRAQRIL		
		201	250
Tr_athaliana	(201) QQRALSNPKIDVIWNSSVVEAYGDGERD-VLGGIKVKNVVTGVDVSDLKVS		
TrxB_sa	(187) QDRAFKNDKIDFIWSHTIKSIN---EKDGKVGSVLTSTKDGEETHEAD		
		251	300
Tr_athaliana	(250) GLFFAIGHEPATK-FLDGGVLDSDGYWVTKPGTTOTSVPGVFAAGDVQD		
TrxB_sa	(234) GVFIYIGMKPLTAPFKDLGITND-VGYIVTKDDMT-TSVPGVFAAGDVRD		
		301	339
Tr_athaliana	(299) KKRYQAITAAGTGCMAALDAEHYLOEIGSQEGKSD---		
TrxB_sa	(282) KGLRQIVTATGDGSTAAQSAEYIEHENDOARSHHHHH		

Figure 16

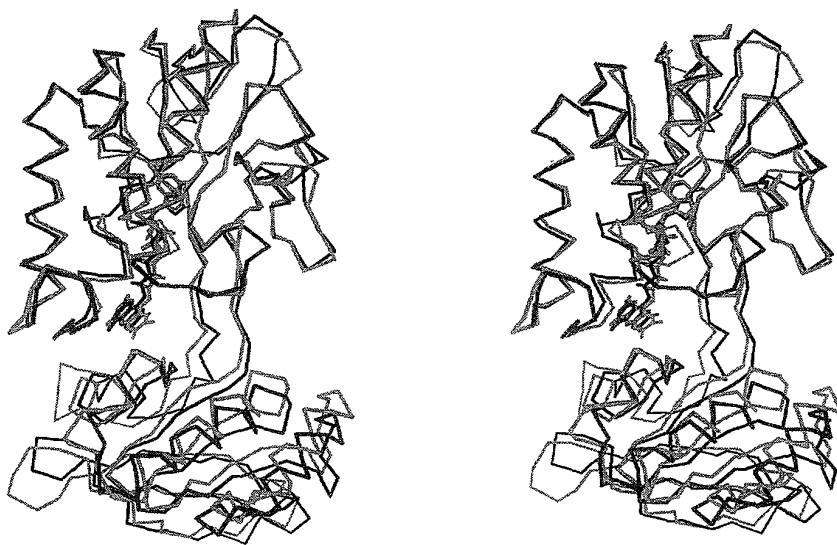


Figure 17

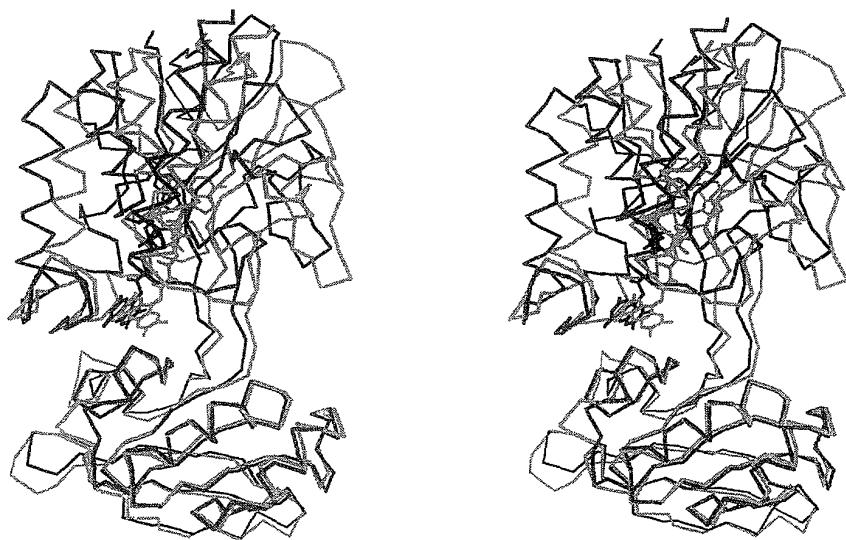


Figure 18

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